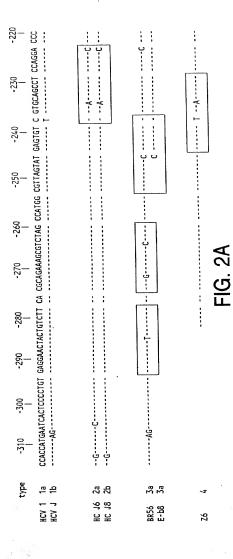
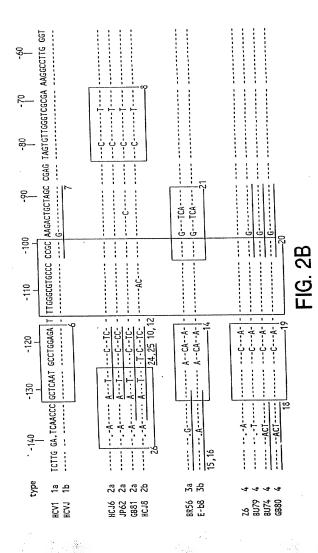


FIG. 1





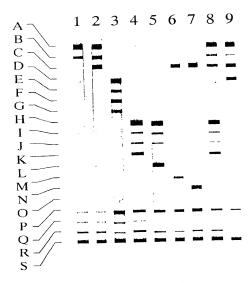


FIG. 3

	ID NO 61 ID NO 62		ID NO 63	SEQ ID NO 64	ID NO 65 ID NO 66		SEQ ID NO 80
	SEQ		SEQ	SEC	SEQ		SEC
-240 -230 -220 -180 -170 -160 -150 -140 ref. agargrechecageccaccageaccaccageaccaccageaccaccageaccaccagaccaccagaccaccagaccaccagaccaccagaccacc		C*			5	6	
-240 -230	E-I	A	-AA				
Type 1a	222	2a	9,9	3c -	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	3b	30 -
Iso- late HCV-1	HCV-J BE82 BE90	HC-J6	HC-J8 BE91	BE92	BR56 BE93 BE94	HCV-TR	BE98

FIG. 4A-1

	SEQ ID NO 67 SEQ ID NO 68 SEQ ID NO 69 SEQ ID NO 70	Q 2	SEQ ID NO 73 SEQ ID NO 74				8 8	SEQ ID NO 72	מבל לבמ	SEQ ID NO 75
Accession -220 -180 -170 -160 -150 -140 ref.		C*CGGT			**	39 40	T*TAIC	51 52		TTTT
-240 -230		TA	TA TA	T	TY		A	A	TY	T
Туре	4444	4p	044 040 0	4C	49	4e	4£	49	4 .	4h
Iso- late	GB48 GB116 GB569 GB358	21	CAM600 CAM736 GB809	24	DK13	GB549	GB438	GB724	BE97	GB487

FIG. 4A-2

	SEQ ID NO 78	SEQ 1D NO 79		
Accession number -140 ref.	TT	A	48 48	CAA
160 -150 +	AA 44 * 45	47		62 T
-170				
-220 -180 -170	* * *	*	C	
40 -230	AA	AA*	A	A
Iso- Type -240	Sa : :	5a	6a	65
Iso-	SA1 BE95	9638	HK1	HK2

FIG. 4A-3

Isolate Type	Туре	-130	-120	-110	-100	06 -	08-	-70	Accession number reference
HCV-1	1a	${\tt ccccrcaatcctggagatttggccgrccccccaagactgctaagactaactaagactagccgaagatagctgcgaagatttgggtcgcgaagaagaagaagaagaagaagaagaagaagaagaag$	CCTGGAGATT	resecerec <u>c</u>	CCCGCAAGAC	rectaeccea 31	GTAGTGTTGC	GTCGCGA	Т
HCV-J BE82 BE90	444		29		ეე-	30			2
HC-J6	2a	ATCTC	DID				T	I	E
HC-J8 BE91	9 9	ATT-CTCAC	T-CTC	AC		-L	TCT	T	4
BE92	2c	ATC	TC	34			C	T	
BRS6 BE93 BE94	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	A - CA -	ACAA ACAC		GTCA GTCA TCA	.A			S.
HCV-TR	33		CA		GTCA	.A			9

FIG. 4B-1

Isolate Type	Туре	-130 +	-120	-110	-100	06-	08-	Acc	Accession number reference
GB48 GB116 GB569 GB358	4 a 4 a 4 a 4 a 4 a 4 a 4 a 4 a 4 a 4 a		CA						
GB549	4.6		CCA	41					
CAM600 CAM736 GB809	40 40 40		-TCA -TCA -TCA	43		C			
GB487	4h		A	5.0					
GB724 BE97	4 g 4 g		A						
SA1 BE96 BE95	ស្រួល		-C	46					7
HK1	6а		1						8
BE98	ЗС .		-CA		g				
GB438	4£ -		CA		g				

FIG. 4B-2

	HG - SA - S	HC-J1	SIVESUIXIEEAIIQUULTAMAYARAL CRITAGAT I INSRENGA INSRENGA INSPENDINGATION OF THE CRITAGATA AND AND AND AND AND AND AND AND AND AN
	No. No.	HCV-H	KK-0XK
		HCV-JK1	K-0I
V-S-A-E-QR	V.S. A-E-QR	HCV-CHINA	IK-Q
V.S. A-E-QR-K-QR-G-R-G-R-G-R-G-R-G-R-G-R-G-R-G-R-G-		HCV-T	IK-0
		HC-14.91	IK-0
	N - V - S - N - E - Q	HCV-TA	IK-Q
N V - S - S - A - E - Q K - Q K - Q	N V - S S - E - Q K - Q K - Q K - Q K - Q K - Q - -	HCV-JT	IK-Q
New Color New	N. V. S. S. R. S. PEE-HI - H. S. C. G. S. R. S. PEE-HI - H. S. R. S. PEE-HI - H. S. R. S. R. S. S. R. S. S. R. S. PEE-HI - H. S. R. S. R. S.	HCV-BK	IK-Q
R. S. RA. S. PEE-HTH. MF - K QT - M. R. S. RA. S. PEE-HTH. MF - K QT - M. R. S. RA. S. PQE - TU'H - M. R. GS - M. S. RA. S. PQE - TU'H - M. R. GS - M. S. RA. S. PEE - T. H. M. R. GS - M. S. R. S. L. S. PEE - T. H. M. R. GT - M. R. R. CT - M. R. R. CT - M. R.	R. S. RA. S. PEE-HTH. MF-K QI R. S. A. S. PQE-TV-H. MI-K-GS R. S. A. S. PQE-TV-H. MI-K-GS R. S. LA. S. PCE-TV-H. MI-K-GS Q. Q. Q. C. E. N. E. E. RV-S C. MF-K-AQ Q. Q. Q. C. E. N. E. E. RV-S C. MY-K-VQ MH-K-QI MH-K-QI K. Q. C. E. R. SA MH-K-QI K. Q. C. E. R. SA MH-K-QI K. Q. C. E. R. SA MH-K-DI K. Q. C. E. R. SA MH-K-DI K. Q. C. C. MH-K-DI MH-M-S. S. S. Q-E-A-R-R-Q-C-MY-K-QQ MH-K-QO MH-M-S MH-M-S MH-M-S MH-K-QO MH-K-QO MH-M-S MH-K-QO MH-K-QO MH-K-QO MH-K-QO MH-K-QO MH-K-QO MH-M-S MH-K-QO MH-M-S MH-K-QO MH-K-QO MH-M-S MH-M-M-S MH-M-M-M-M-M-M-M-M-M-M-M-M-M-M-M-M-M-M-	BE90	IK-Q
R. S. A. S. PQE - TV + H - W - K - QS - W - K - QS - W - S - PQE - TV + H - W - W - K - QS - W - S - PQE - TV + H - W - W - W - W - W - W - W - W - W -	R. S. S. A. S. PQE - TV H	HC-16	MFK-QT
R. S A. S-PQE - TV-H MI K-QS MI K-QT MI MI	R. S A. S. PQE - TV'H M K-QS	HC-J8	WK-0S
		BE91	MIK-QS
		BE92	MLK-QT
		Π	CMFK-AQ
		17	CMYK-VQ
	H E E. E K SA I MY - K. LQ	BE93	CMFK-AQ
		19	IWYK-LQ
KVEVE-EKTAMHK-DLKV-EVE-ERTAMHDLKV-EVE-EKTAMHK-DLKV-EVE-EKTAMHK-DLRV-EVE-EKV-SAMHK-DLRKVSAMHK-DLRKVSAMHK-DLRKVSA	Kvevevee-k-ra-namh-k-dl Kvevee-ee-k-ra-namh-k-dl Kvevee-e-k-ra	110	IMYK-LQ
KvEvE-ERTAMHDLKvEvE-EKTAMHKDLRvEvE-EKTAMHK-DLRvEvE-EKV-SAMHK-DLRvEvE-EKV-AAMHK-DLRvEvE-EKV-AA		GB48	MHK-DL
KVEVE-EKVTAMHK-DLRV-EVE-E-KVSAMHK-DLRV-EVE-E-KVSAMHK-DLRVVEVE-E-KVAAMHK-DLRVVAAMHK-DLRVVAAMHK-DLRVVAAMHK-DL		GB116	TGHW
KVEVE-E-K-1ANH-K-DLRE-E-KV-SANY-K-DLQR-KV-EVE-E-KV-AANH-K-DLH-MSSQ-E-A-R-R-Q	KvEvE-EKTAMHK-DL RXEKV-SAMHK-DLG RXEKV-AAMHK-DL HMSSGKRGCMYK-QQ FIG. 5A	GB215	MHK-DL
REE-E-KV-SA	R	GB358	MHK-DL
R-KVEVE-E-KV-4A	R-KV-EVE-E-KV-AA	GB549	
H-MSSQ-EARQCMYK-QQ	H-M-08-08-08-6-A-R-08-08-08-04-48-00-0-0	GB809	
HMSSLY-Q-ERQCMYK-QQ	FIG. 5A	BE95	CMYK-QQ
	FIG.	CHR18	

	101	
HCV-1	SAGVQEDAASLRA	
HC-J1		
HCV-H		
HCV-J	TA	
HCV-JK1	TV	
HCV-CHINA	TV	
HCV-T	TV	
HC-J4.91	TA	
HCV-TA	TV	
HCV-JT	TV	
HCV-BK ·	TV	
BE90	TV	82
HC-J6	-Q-TEERN	
HC-J8	-Q-NEERN	
BE91	-Q-NEERN	83
BE92	-Q-TEERN	84
T1	-DDR-A	
T7	-DDRTA	
BE93	-DDR-A	85
T9	-C~-ER-A	
T10	-CER-A	
GB48	-DEKRP-G-	86
GB116	-DEKRA-G-	87
GB215	-DEKRA-GV	88
GB358	-DEKRA-G-	89
GB549	-GERA	90
GB809	-GEKRA-G-	91
BE95	-Q-THE	92
CHR18	-Q-THK	

FIG. 5B



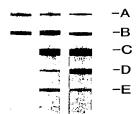


FIG. 6

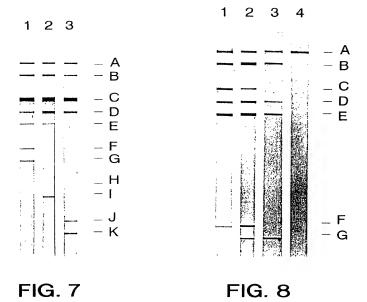


FIG. 9

FIG. 10